

Table 1. Amino acid sequence of peptides hydrolyzed by human glandular kallikrein 2 (HK2)

| | | | | | | |
|---|---|---|---|---|---|-----------------|
| G | K | A | R | A | F | (SEQ. ID NO: 1) |
| G | K | A | V | R | Q | (SEQ ID NO: 2) |
| G | K | A | Y | F | M | (SEQ ID NO: 3) |
| G | K | A | E | K | V | (SEQ ID NO: 4) |
| G | K | A | F | R | K | (SEQ ID NO: 5) |
| G | K | A | K | P | R | (SEQ ID NO: 6) |
| G | K | A | A | Y | Y | (SEQ ID NO: 7) |
| G | K | A | M | Y | H | (SEQ ID NO: 8) |
| G | K | A | F | R | R | (SEQ ID NO: 9) |
| G | K | A | L | Q | R | (SEQ ID NO: 10) |
| G | K | A | M | R | Q | (SEQ ID NO: 11) |
| G | K | A | A | L | M | (SEQ ID NO: 12) |
| G | K | A | Q | G | F | (SEQ ID NO: 13) |
| G | K | A | N | M | N | (SEQ ID NO: 14) |

Random library constructed with sequence

NO₂-Y-G-K-A-X₁-X₂-X₃-Dap-F-K(ABZ)

Where NO₂-Y is nitrotyrosine quencher

X₁, X₂, X₃ are random amino acids consisting

of all natural L-amino acids except cysteine (n=19)

Dap is diaminopropanoate; K(ABZ) is lysine coupled to

fluorophore-aminobenzoyl acid (ABZ)

HK2 cleavage sites denoted by single or double //

FIGURE 1

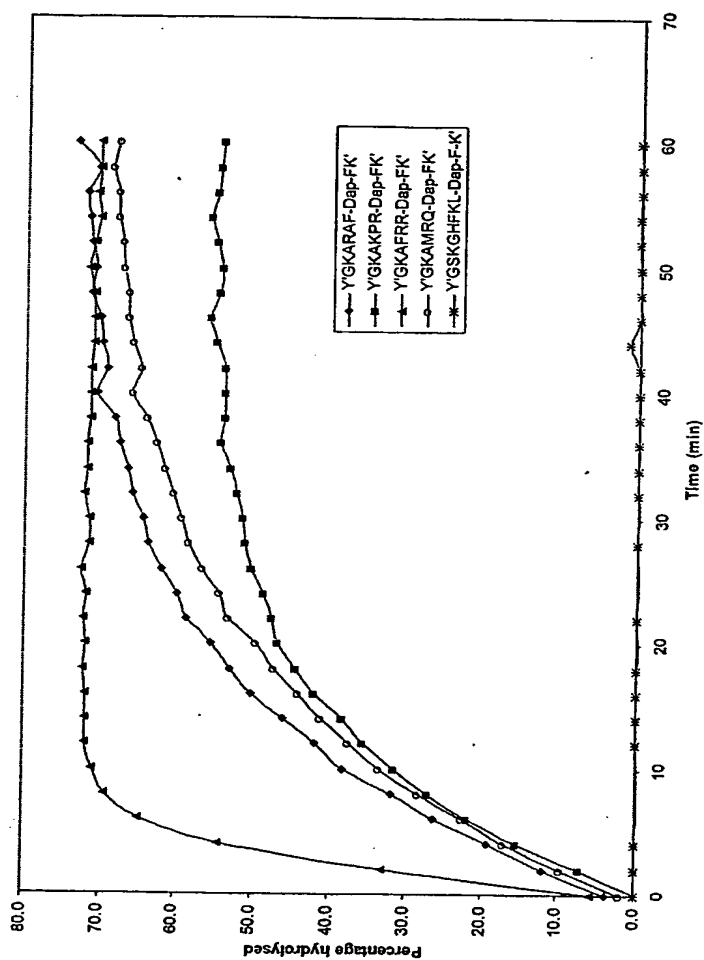


Figure 2

hK2 Hydrolysis Sites

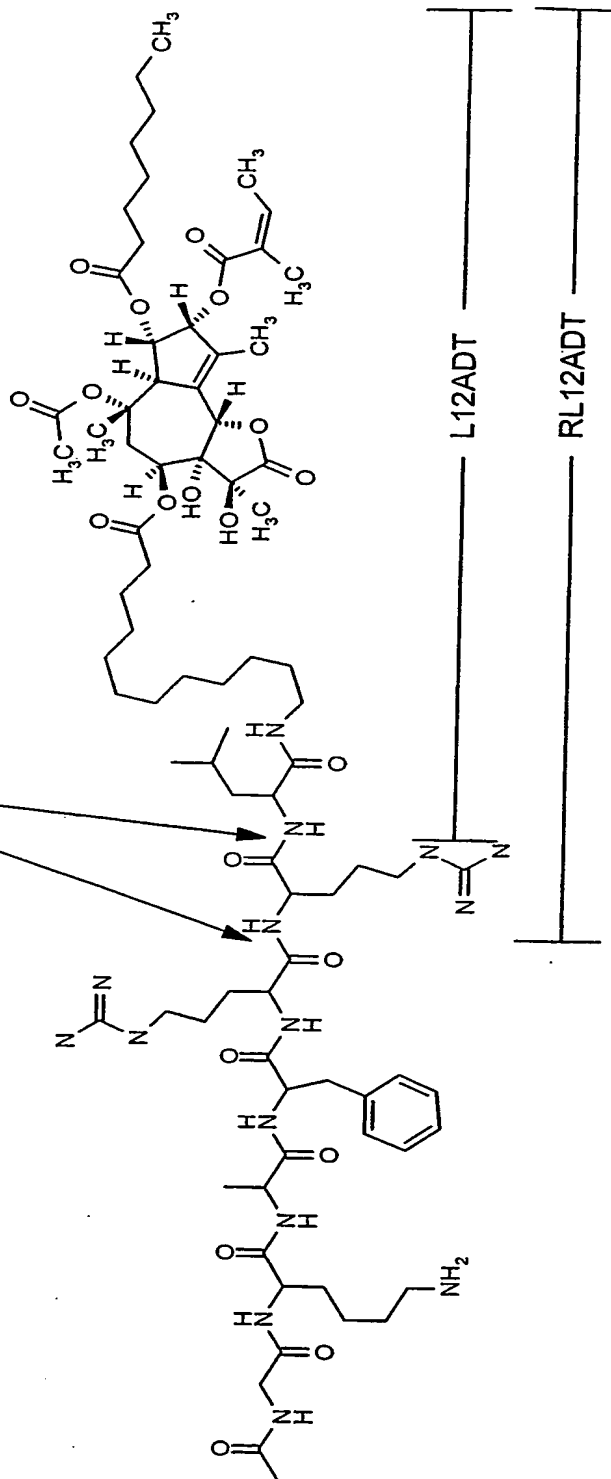
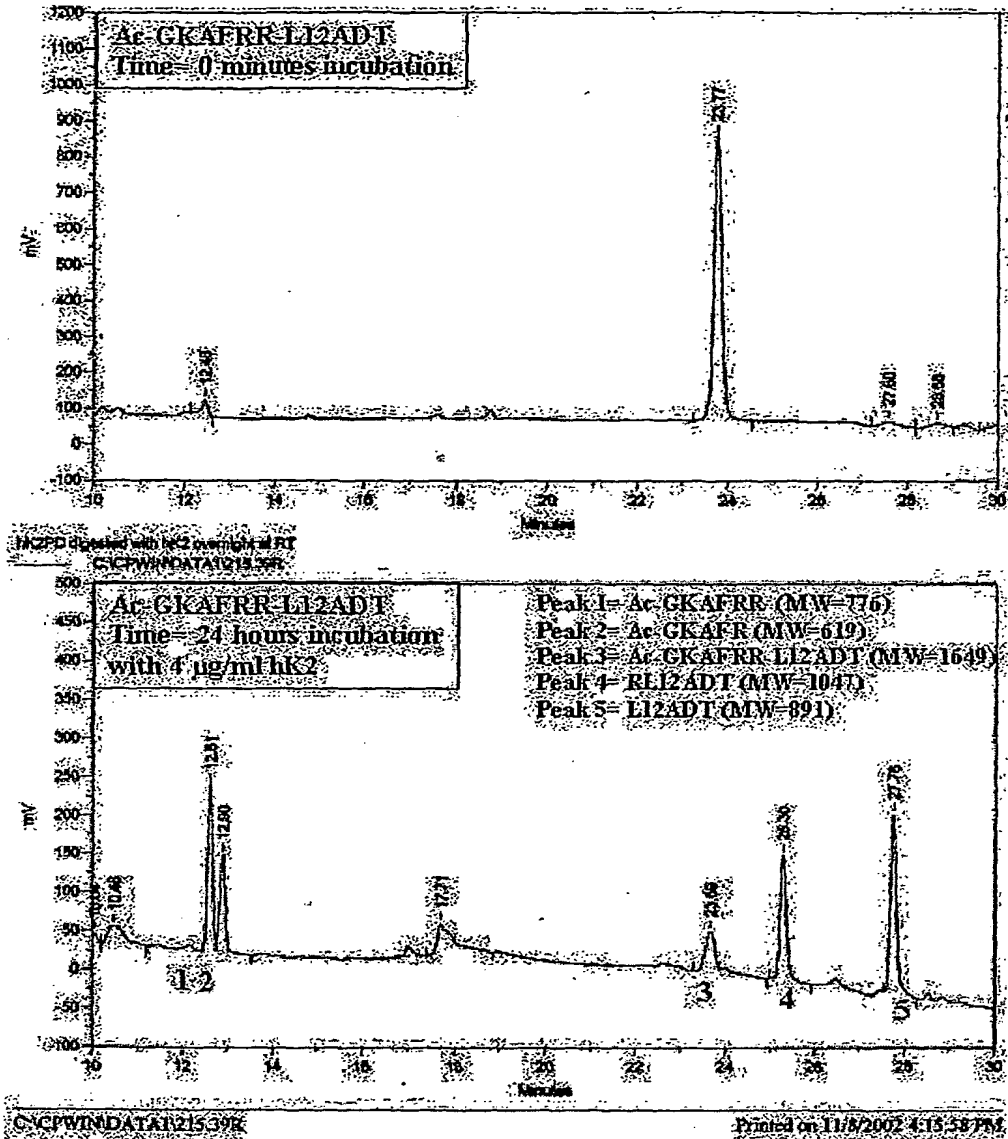
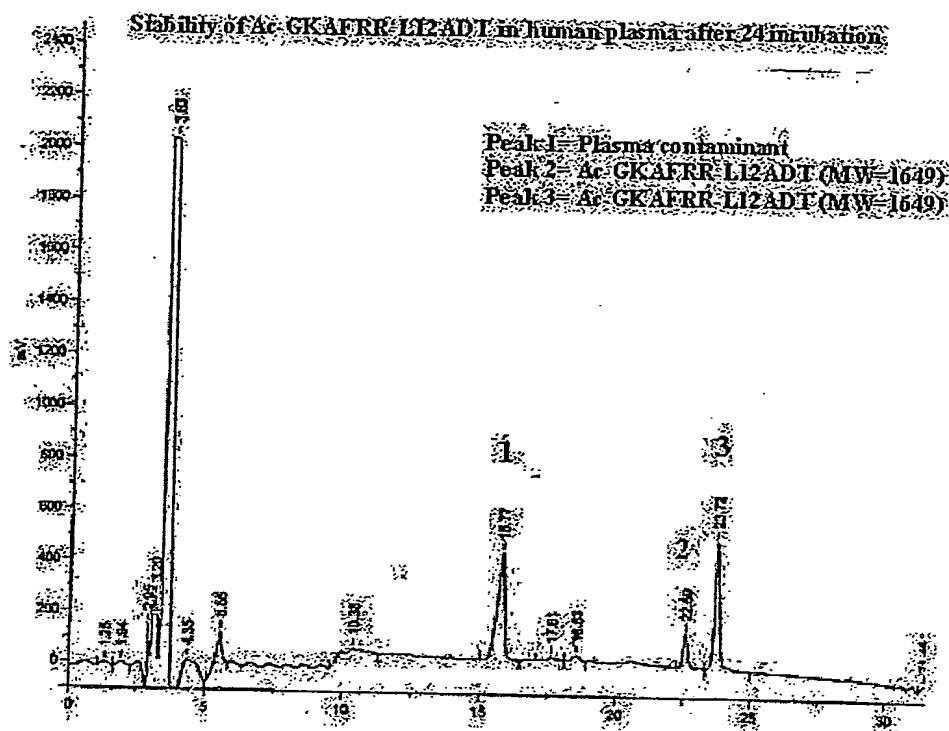


FIGURE 3



HPLC analysis of hydrolysis of hK2 prodrug Ac-GKAFRR-LI2ADT by hK2 (4 µg/ml) over 24 hr. incubated in 50 mM TRIS, 0.1 M NaCl, pH 7.3 at room temperature. Mass of each peak confirmed by MALDI-TOF mass spectrometric analysis (see figure 3 for mass profiles).

FIGURE 4



HPLC analysis of Ac-GKAFRR-L12ADT incubated in 50% human plasma for 24 hrs at room temperature. Peak 1 represents unidentified plasma contaminant that was also present in control plasma. Peak 2 and 3 both represent Ac-GKAFRR-L12ADT as confirmed by MALDI-TOF mass spectrometric analysis.

FIGURE 5

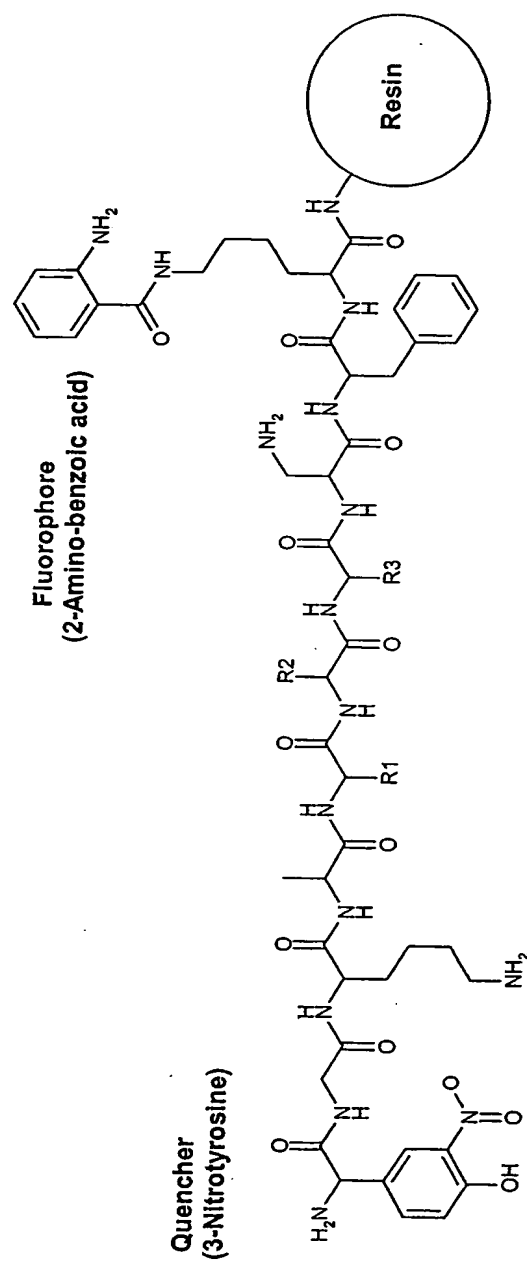


Figure 6

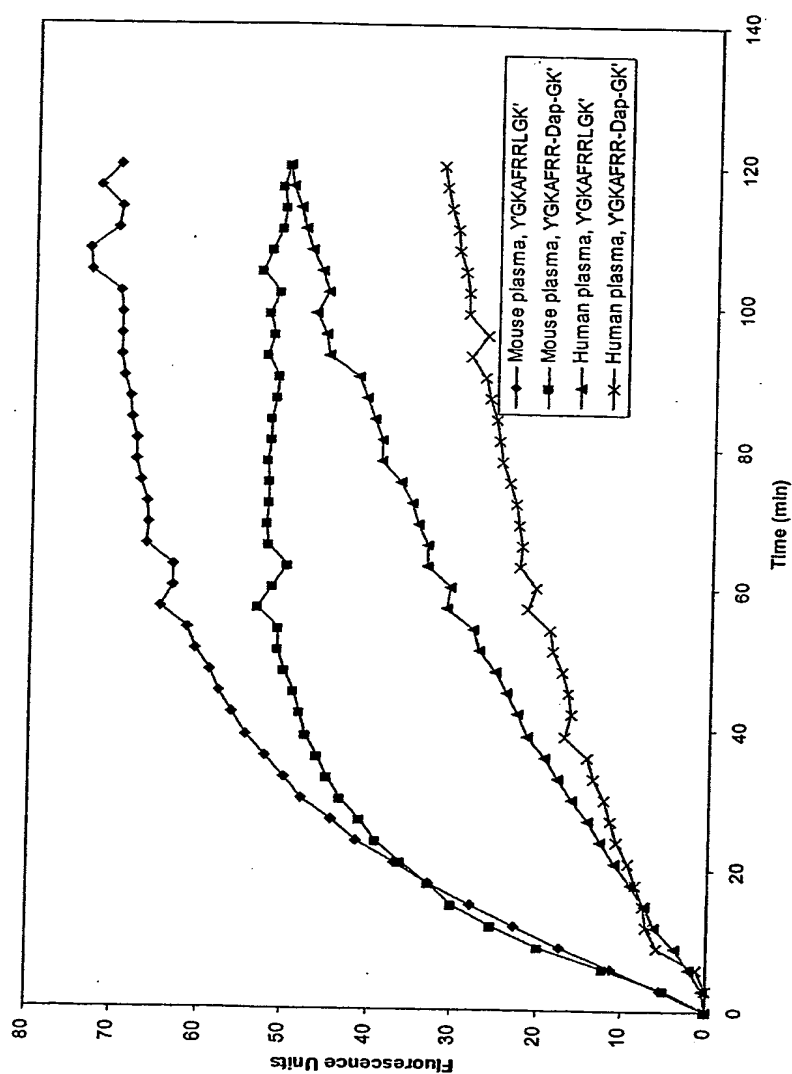


FIGURE 7

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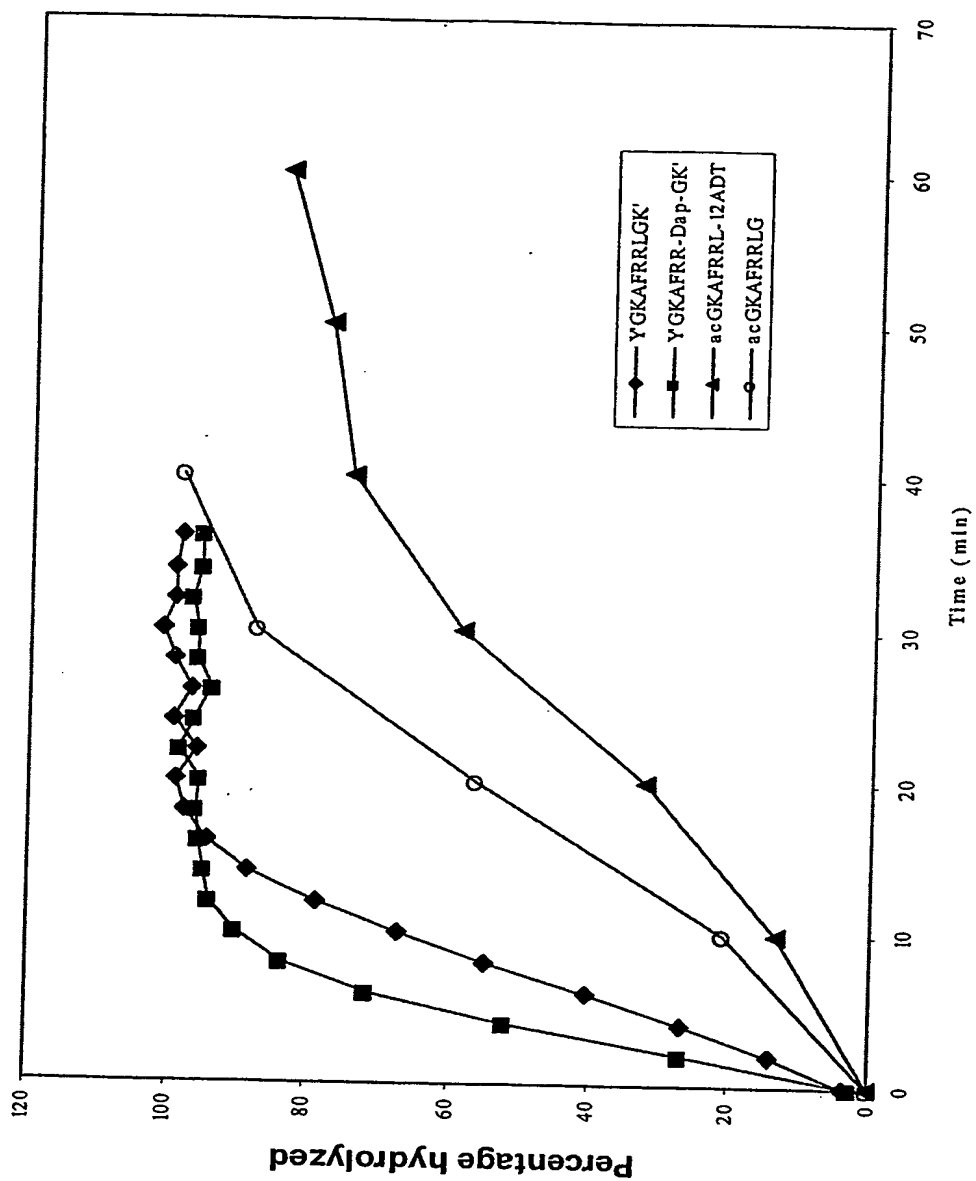


FIGURE 8

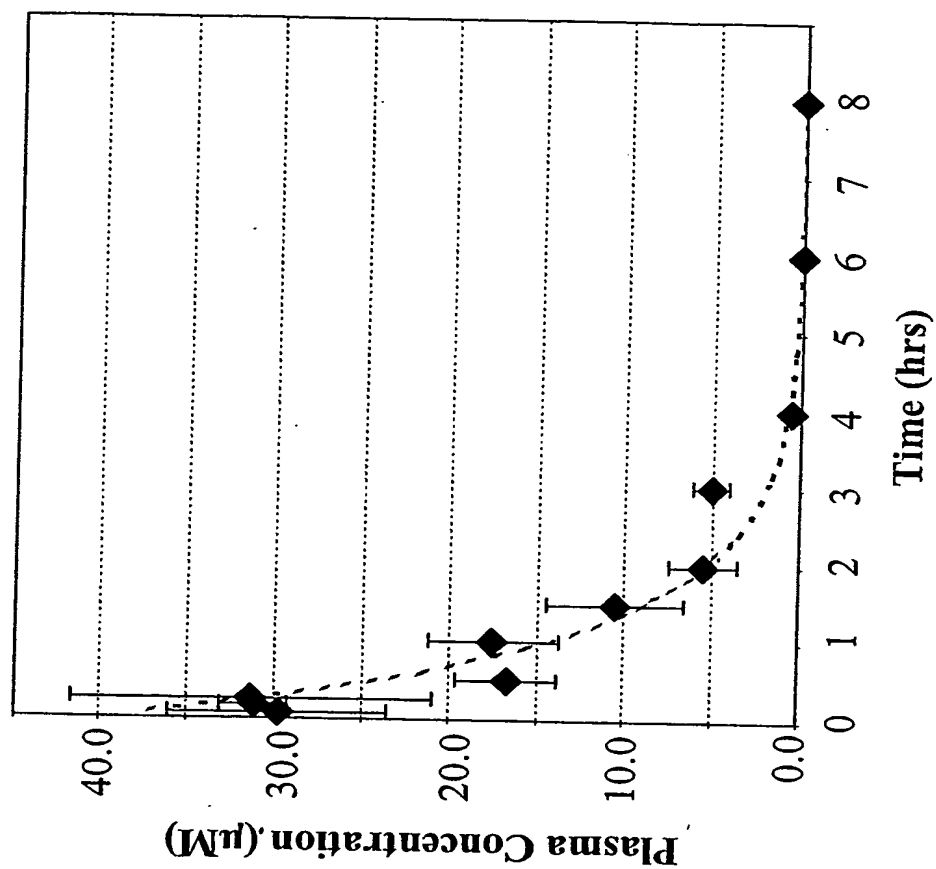


FIGURE 9